## STIC-Biotech/ChemLib

From: Sent:

To: Subject: Gibbs, Terra Friday, September 13, 2002 1:41 PM STIC-Biotech/ChemLib

Sequence Search

Could you please search SEQ ID NO: 3 of serial number 10/003354

Please do a length limited search of 50 nucleotides or less. Also no EST's.

Terra Gibbs #79523 **AU 1635** Mailbox 11E12 306-3221

THANK YOU!

**Edward Hart** Technicai Info. Specialist STIC/Biotech CMI 6B02 Tel: 305-9203

Searcher:
Phone:
Location:
Date Picked Up: 41,19402
Date Completed: (1/8)(0)
Searcher Prep/Review:
Clerical:
Online time:

TYPE OF SEARCH: ,
NA Sequences:/
AA Sequences:
Structures:
Bibliographic:
Litigation:
Full text:
Patent Family:
Other:

VENDOR/COST (v	here applic.)
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## ALIGNMENTS

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DEFINITION
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A59476
A59476.1
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Chamberlain,S., Pook,M.A., Doudney,C.,
Garcia-Valdecasas,J.J. and C.
GENE FOR FRIEDREICH'S ATAXIA
Patent: WO 9705234-A 26 13-FEB-1997;
IMPERIAL COLLEGE (GB)
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26 from Patent W09705234.
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                                                                                                                                   Shimkets,R.A. and Leach,M.
Nucleic acids containing single nucleotide
methods of use thereof
Patent: WO 0140521-A 102 07-JUN-2001;
Curagen Corporation (US)
                                                                                                                                                                                                                                                                       AX156774
Sequence 102 from Patent
AX156774
AX156774.1 GI:14538105
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Chamberlain, S., Pook, M.A.,
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       2256 cagagagttgaggaccgcagcatcccctccactccagagttgg 2298
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Artificial sequences.
JP 1995194400-A/11
01-AUG-1995
21-OCT-1994 JP 1994282915
22-OCT-1993 JP 93P 265144
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SUZUKI KATSUHIRO
C12Q1/68,G01N33/50//C12N15/09;
CAGACTATTGAAGTCGGCCTCATCCCCACCACTGCATACATGG
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AX159492
Sequence
AX159492
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Immunostimulatory rna/dna hybrid molecules
Patent: WO 0193902-A 349 13-DEC-2001;
Biosynexus Incorporated (US)
Location/Qualifiers
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Patent: WO
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Schweizer,M.
BIOLOGICAL ASSAY FOR TESTING THE CARCINOGENIC PROPERTIES
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Sequence 10 from Patent WO9845476
A84538
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synthetic construct
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FOOD RESEARCH (GB); SCHWEIZER
Location/Qualifiers
             2820
                                                                                                                                                                                     /organism="synthetic construct"
/db_xref="taxon:32630"
/note="Synthetic HDR"
/note="Synthetic HDR"
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/db_xref="taxon:32644"
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from Patent WO0140521.
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Pred. No. 1.5e
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Pred. No. 1.5e+06;
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                                               Query Match 0.6%;
Best Local Similarity 76.5%;
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 50)
Shimkets,R.A. and Leach,M.
Nucleic acids containing single nucleotide polymorphi
methods of use thereof
Patent: WO 0140521-A 2822 07-JUN-2001;
Curagen Corporation (US)
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae;
1 (bases 1 to 50)
Shimkets,R.A. and Leach,M.
Nucleic acids containing single nucleotide polymorphi
methods of use thereof
Patent: WO 0140521-A 2820 07-JUN-2001;
Curagen Corporation (US)
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/db_xref="taxon:9606"
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/note="Nucleotide deleted between
Accession number cg42513366"
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/db_xref="taxon:9606"
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AR069130
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AX159496
Sequence 1 from Patent WO0144811.
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Patent: WO 0140521-A 2824 07-JUN-2001;
Curagen Corporation (US)
                                                                                                                                                                                                                            Vaidya,T., Bakhiet,A., Olsson,T., Kristensson,K.
Lymphocyte stimulating factor
Patent: US 5891439-A 14 06-APR-1999;
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/db_xref="taxon:9606"
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Accession number cg42513366"
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Sequence 47:
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AX161426.1
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I32124
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Singh, S., Pease, J.S., Sadakian, J., Wagner, D.B. and Ullman, E. Light-emitting particles for diagnostic and therapeutic use Patent: WO 0.144811-A 1 21-JUN-2001;
DADE BEHRING INC. (US)
            Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Bouma, S.R., Khalil, O.S.
Method for detection of
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AX172348.1
 Shimkets, R.A.
                                                                                                                                                                                                                                                                                                                                                                                                Unknown
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                                                                                                                                                                                                                                                                                                                      Patent: US 5585242-A 14 17-DEC-1996;
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/db_xref="taxon:32630"
/note="Oligonucleotide Probe"
3 c 6 g 25 t
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7 c 3 g
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          1 to 50)
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                                                                                               Patent
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Pred. No. 2e+0
0; Mismatches
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Pred. No. 2.2e+06;
0; Mismatches 7;
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nucleic acid using
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WO0140521.
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5 5585242.
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2e+06;
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REFERENCE
AUTHORS
TITLE
VERSION
KEYWORDS
SOURCE
                                     RESULT 15
AX040470/c
LOCUS
DEFINITION
ACCESSION
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AX199596/c
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Best Local Similarity
Matches 25; Conserv
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Best Local Similarity 64.0
Matches 31; Conservative
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                                                                                                                                               2307 gattttcaactggccaaccctttgcctccactattgaatttttttcag 2354
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Patent:
Curagen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        50 bp
Sequence 526 from Patent W00151670.
AX199596 GI:15390030
                       AX040470 41 bp
Sequence 10 from Patent W00063365.
AX040470 GI:11230262
                                                                                                                                                                                                                                                                                                                                                                                              methods of use thereof
Patent: WO 0151670-A 526 19-JUL-2001,
Curagen Corporation (US)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 50)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
synthetic construct.
                                                                                                                                                                                                                                                                                                                                                                                                                                      Shimkets,R.A. and Leach,M.D.
Nucleic acids containing single nucleotide polymorphisms and
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/db_xref="taxon:9606"
25...26
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/db_xref="taxon:9606"
25..26
                                                                                                                                                                                                                                                                                         /note="Nucleotide deleted between Accession number cg42855114" 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="Nucleotide deleted between bases Accession number cg43969665" 26
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Pred. No. 2.3e
0; Mismatches
                                                                                                                                                                                  Score 20.8; DB 6;
Pred. No. 2.3e+06;
0; Mismatches 17;
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2.3e+06;
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TITLE
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                                                                                   3566 ggagggctgggtcttaggggaaaggaatggggaagc 3600
                                                                       36
                                                                     GGTGGGTGGGGTATTAGGGGAGGGAGGGAAGC 2
                                                                                                                                                                                                                                                                                               artificial sequence.

1 (bases 1 to 41)
Belotserkovski, B., Reddy, G. and Zarling,
Locked nucleic acid hybrids and methods o
Patent: WO 0063365-A 10 26-OCT-2000;
Pangene Corporation (US)
                                                                                                                                                                                                                                                                                                                                                                                      synthetic construct
                                                                                                                                        Similarity
                                                                                                                            Conservative
                                                                                                                                                                                                             /organism="synthetic construct"
/db_xref="taxon:32630"
/note="Triplex forming DNA"
25 c 2 g 7 t
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              2002, 01:15:45
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                                                                                                                           Score 20.6; DB 6; Pred. No. 2.6e+06; 0; Mismatches 9
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Post-processing: Minimum Match 0% Maximum Match 100%
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Maximum DB seq
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222.22.66
222.22.66
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Gapop 10.0 , Gapext 1.0
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Copyright (c) 1993 - 2000 Comp
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Human SNP oligonuc
Human SNP oligonuc
(Ser4Gly)2 linker
DNA encoding (Ser4
Human silent SNP c
HHV6 virus p41 gen
Human SNP oligonuc
IgE receptor beta
                                                                                                                                                                                                                                                                                                                      Description
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22 22 22 22 22 22 23 24 24 24 25 26 27 27 27 27 27 27 27 27 27 27 27 27 27	16 22 22 22 22 22 22
AAIT5879 AAIT5881 AAIT5883 AAIL28737 AAZ211524 AAZ11152 AAZ1169 AAZ67568 AAZ67568 AAIC28914 AAZ17813 AAIH9745 AAZ23570 AAC69633 AAIH9745 AAC69633 AAIH9749 AAC69633 AAH97394 AAIL29036 AAL29036 AAL29036 AAL29036 AAL29036 AAL29036 AAL29036 AAL29036 AAL3957659 AAL29036 AAL3957659 AAL3967659 AAL3967659 AAL3967659 AAL3967659 AAL3967659 AAL3967659 AAL3967659 AAL3967659 AAL3967659 AAL397894 AAL3967659 AAL3967669 AAL3967669 AAL3967669 AAL3967669 AAL39676768	AAT04081 AAL34072 AAL28913 AAH20339 AAL31293 AAV83644 AAL30217 AAL27921 AAL28580
silen silen silen silen silen silen for H 1 us map-r silen silen codi ion se phila phila codi SNP o	Trypsin inhibitory Human SNP oligonuc Human SNP oligonuc HHV6 virus p41 gen Human SNP oligonuc Oligonucleotide us Human SNP oligonuc Human SNP oligonuc Human SNP oligonuc

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## ALIGNMENTS

RESULT

AAT84374 standard; DNA;

49

ВP

12-NOV-1997 (first entry)

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Chamberlain S, Hillermann R,
                                                                                                                                                               STM7.I gene; Friedreich's ataxia; FRDA; neurodegeneration; phosphatidylinositol-4-phosphate 5-kinase; PtdInsP5K; transgenic animal; animal model; diagnosis; therapy; STM7.IIIa; STM7.IIIb; STM7.IIIC; ss.
WPI; 1997-145674/13
                                                               28-JUL-1995;
26-JUL-1995;
                                                                                                                                                                                                              Friedreich's ataxia STM7 gene exon 11.
                                                                                         24-JUL-1996;
                                                                                                            13-FEB-1997.
                                                                                                                             WO9705234-A2
                                            (UNLO ) IMPERIAL COLLEGE SCI TECHNOLOGY & MED
                                                                                                                                                Homo sapiens
                   Pook MA;
                                                               95GB-0015508
95GB-0015309
                                                                                          96WO-GB01786
                           Doudney
                           CWE,
                            Garcia-Valdecasas JJC;
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RESULT 2
AAL30887/c
В
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Best Local S
Matches 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA sequences (AAT84364-86) are provided for exons 1-23, respectively, of the human STM7 gene associated with Friedreich's ataxia (FRDA). The STM7.I gene (see AAT84358) comprises exons 1-16 of the gene, while splice variant STM7.IIIa includes exons 1-13, 17 and 19-22, splice variant STM7.IIIb includes exons 1-15, 17, 19, 21-22 and splice variant STM7.IIIc includes exons 1-15, 17, 19, 22-22 and splice variant STM7.IIIc includes exons 1-15 and 19-22. STM7 nucleic acids can be used to produce STM7 polypeptides (see AAM00978) in transfected host cells, to design probes (esp. oligonucleotides from exons 1-16, partic. 7-11) used in a claimed method for determining an inherited predisposition to FRDA, to generate transgenic animal models of FRDA and in somatic cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New gene, STM7, and new protein causing Friedreich's Ataxia encodes phosphatidylinositol phosphate kinase isoform; also splivariants and transgenic animals, for diagnosing pre-disposition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1591 gtttgttaagaagttggagcactcttggaaagccctggtacatgacgg 1638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Friedreich's ataxia
                                                                                                                                                                                                                                                                                                                                                                                                     neuroprotective; antimicrobial; gene therapy; vaccine; amylase; camyloid protein; angiopoietin; apoptosis related protein; cadhering cyclin; polymerase; oncogene; histone; kinase; colony stimulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAL30887
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 49 BP; 15 A; 5 C; 13 G; 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human SNP oligonucleotide #4095
                     polymorphic nucleic acids encoding e.g. amylases, cyclins, poloncogenes and histones, useful for diagnosing and treating, e cancer, autoimmune diseases and infections.
                                                                                                                                                                      28-DEC-1999;
27-DEC-2000;
                                                                                                                                                                                                                                                  05-JUL-2001
                                                                                                                                                                                                                                                                                 WO200147944-A2
                                                                                                                                                                                                                                                                                                                                                            multifactorial
                                                                                                                                                                                                                                                                                                                                                                       complement related protein; cytochrome; kinesin; cytok interleukin; G-protein coupled receptor; thioesterase;
                cancer,
                                                                            WPI; 2001-465210/50.
                                                                                                                                                                                                                    28-DEC-2000;
                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 gttaatgaagaagttagaacattcctggaaagctcttgtttatgatgg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      l Similarity
36; Conser
                                                                                                                                                                                                                                                                                                                                           system
              autoimmune diseases and infections
                                                                                                                                         CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fig 16; 82pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard;
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nilarity 75.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                        2000US-0173419
                                                                                                                                                                                                                      2000WO-US35498
                                                                                                           Leach M;
                                                                                                                                                                                       99US-0173419.
                                                                                                                                                                                                                                                                                                                                             disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                           autoimmune disease; ss.
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Pred.
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No. 4e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12;
                                                                                                                                                                                                                                                                                                                                                                                         cytokine; interferon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length
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                                             cyclins, polymerases,
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                                                                                                                                                                                                                                                                                                                                                                           inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       49;
                                                                                                                                                                                                                                                                                                                                                                                                                          cadherin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                         cancer;
                                                                                                                                                                                                                                                                                                                                                                                                           factor;
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RESULT
AAL31735
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                dd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                    neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cano amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating fa complement related protein; cytochrome; kinesin; cytokine; interfero interleukin; G-protein coupled receptor; thioesterase; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2577 ttttttttttctgaaaaaaggaaaaagcacacagcacacaattt 2620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; Page 2562; 4143pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human SNP oligonucleotide #4943.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             24-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAL31735;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 50
                                                                                                                                                                                          28-DEC-1999; 99US-0173419.
27-DEC-2000; 2000US-0173419.
                                                                                                                                                                                                                                                                                               WO200147944-A2
                                                                                                                                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                           nervous
                                                                                                                                                                                                                                                                                                                                                                         multifactorial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAL31735 standard;
                                                Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and cancer authorismune diseases and infections.
                                                                                                                                                                                                                                      28-DEC-2000; 2000WO-US35498
                                                                                                                                                                                                                                                                    05-JUL-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic;
                                                                                                                                                               (CURA-) CURAGEN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TCTTTTTTTAAAAAAAAAAGCAAAAAAGCCATACAACACATATT 1
                                                                                                     2001-465210/50
                                          autoimmune diseases
                                                                                                                                   RΑ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 16 A; 3 C; 6 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                   Leach
                                                                                                                                                                                                                                                                                                                                                           disease;
disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%;
                                                                                                                                                                                                                                                                                                                                                            autoimmune disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВP
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                                          and infections
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ή.
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No. 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22;
                                                          and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                           cyclins, polymerases treating, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          50;
                                                                                                                                                                                                                                                                                                                                                                                                          interferon;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                        factor;
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Claim 1;

Page

2809; 4143pp;

English

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당.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           To variants of proteins related to amylases, amyloid proteins, angiopoietin, angiopoietin, sepoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, chistones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, complement coupled receptors and thioesterases. The present sequence is come such oligonucleotide. The oligonucleotides and the peptides encoded come such oligonucleotide. The oligonucleotides and treatment of come such oligonucleotide with inappropriate expression of the proteins listed by them may be used in the prevented, diagnosis and treatment of companies associated with inappropriate expression of the proteins listed convertion. Disorders that may be prevented, diagnosed and/or treated include convertionable convertion and convertion and convertion of systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, cancer checkenia), diseases of the nervous system and an infection of pathogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
          AAQ99068 is a (Ser4Gly)2 substrate linker which is used in a fibrobly growth factor (FGF)/saporin fusion protein (FP) to increase FP flexibility. DNA sequences encoding FGF/saporin fusion proteins are useful for targetting saporin to a cell carrying the FGF receptor. Sequences used to link together FGFs and other sequences which may be beneficially targetted to a cell (targetted agents, TA) may be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ99068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2566 ctttcttcttttttttttttttgaaaaaag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention relates to oligonucleotides variants of proteins related to amylases, amyloid
                                                                                                                                        Claim
                                                                                                                                                                                                       New conjugates of growth factor receptor ligand and targetted agent
                                                                                                                                                                                                                                                                            Baird
                                                                                                                                                                                                                                                                                                            (PRIZ-) PRIZM
                                                                                                                                                                                                                                                                                                                                              15-MAR-1994;
15-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                15-MAR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                 21-SEP-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conjugate; fibroblast growth factor; FGF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-MAY-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (Ser4Gly)2 linker used in FGF/saporin fusion
                                                                                                                                                                                        partic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3 ctttattattatttttttttttgaaaaaag 33
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                                                                                                                                                                                                                                        1995-336820/43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     proliferation;
                                                                                                                                        ن.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                l Similarity
27; Conserv
 by a sequence
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                                                                                                                                       Page 110;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard;
                                                                                                                                                                     DNA or cytotoxin, used to control >.g. to prevent growth of pterygii
                                                                                                                                                                                                                                                                          Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP;
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                                                                                                                                                                                                                                                                                                            PHARM INC
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94US-0213446
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                                                                                                                                     204pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     regulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.7%;
 which can increase serum stability
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   æ
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                                                                                                                                                                                                                                                                            Nova
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Pred. No. 6.3e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  pterygia;
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                                                                                                                                                                                                                                                                         Sosnowski
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   corneal clouding;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cytotoxin; saporin; eye;
                                                                                                                                                                     cell proliferation in and corneal clouding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein,
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                                                                                                   fibroblast
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RESULT
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Best Local
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targetted delivery of cytotoxins or produg-converting enzymes proliferating cells. The gene delivery composition comprises a that binds the fibroblast growth factor receptor (FGFR) which i or chemically conjugated to a nucleic acid hinding in the conjugated to a nucleic acid hinding is a conjugated to a nucleic acid hindin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-MAR-1994;
15-MAR-1994;
29-AUG-1994;
13-SEP-1994;
16-MAY-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       intracellular availability of the TA. The conjugates of the invention are used to inhibit cell proliferation in cells carrying the particular growth factor receptor; also when TA is DNA it can be used to deliver this to cells (for gene therapy). A specific application is to prevent excessive proliferation of epithelial cells, fibroblasts and keratinocytes in the anterior eye after surgery, partic. to prevent recurrence of pterygii after surgical removal, closure of trabeculectomy after glaucoma surgery and corneal clouding after excimer laser treatment. Other conditions which may be treated include tumours, restenosis, psoriasis, Dupuytren's contracture, diabetic complications, Kaposi's sarcoma and rheumatoid arthritis.
                                                                                                                                                                                                                          Gene delivery system, useful for treating or preventing cancer and rheumatoid arthritis, comprises receptor-internalized ligand linked nucleic acid binding domain and nucleic acid -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FGFR-binding protein; nucleic acid binding protein; receptor-internalised ligand; cytotoxin; saporin; gene therapy; cytocide; antiproliferative; cancer; melanoma; diabetic retinopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rheumatoid arthritis; restenosis, Dupuytren's contracture; eczema; nuclear translocation signal; NTS; cytoplasmic translocation signal; endosome-disruptive pept
                                                                                                                                                                               Claim
                                                                                                                                                                                                                                                                                                                                                                                       Chandler LA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Targetted gene delivery; fibroblast growth factor
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                                                                                                                                                                                                                                                                                                                                                                                                                                          (SELE-) SELECTIVE GENETICS INC.
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                                                                                                                                                                               Column 33;
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                                                                                                                                                                                                                                                                                                                                                                                       Sosnowski BA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94US-0213446.
94US-0213447.
94US-0297961.
94US-0305771.
95US-0441979.
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75.7%;
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Pred. No. 2.2e+04
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RESULT
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ID AAI7
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Matches 28
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                                                                                                                                                                                                                                                                                                                                                                                                                Human; single nucleotide polymorphism; SNP; genome; gene therapy; protein therapy; vaccine; probe; diagnostic assay; detection; quantitation; restorative therapy; polymorphic; ds.
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                                                                                                                                                                                                                                                                                 30-NOV-2000; 2000WO-US32758
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                                             Claim 1; Page 85; 2653pp; English
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29-NOV-2000;
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                                                                                            Polymorphic nucleic acid sequences,
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AAI73060 to AAI79867 represent isolated human polymorphic polynucleotide sequences (I), which contain single nucleotide polymorphisms (SNPs).

This invention relates to a method for detecting the presence of a specific nucleic acid in a sample containing DNA. The method comprises performing solid phase amplification of DNA template (SPADT). 5' and 3' primers are irreversible bound to a solid support, and the DNA from a sample is absorbed and reversibly bound, incubated under amplification reaction conditions and the presence of the specific target DNA is

Column

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RESULT
AAH20340
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAM53114 to AAM53329 represent peptides related to human polymorphic polynucleotide sequences. The sequences can be used in gene and protein therapy, and in vaccine production. (I) and the polypeptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of polymorphic polypeptides. For example, (I) may be used to treat disorders by rectifying mutations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antibodies may also be used to down regulate expression and activity the antibodies may also be used as diagnostic agents for detecting presence of polymorphic polypeptides in samples.
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                                                                                                                      Detecting the presence of a specific nucleic acid in a sample containing DNA, useful in scanning large genomic fragments for presence of genes or gene families, comprises performing solid amplification of DNA template
                                                                                                                                                                                                                                                                                                                                                                                                                               primer; solid phase amplification of DNA template;
genomic scanning; bacterial diagnostic; p41; HHV6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          detected. The method is useful for detecting the presence of a specific nucleic acid (e.g. bacterial, viral or parasitic NNA) in a sample or in a cell. SPADT may be used for scanning large genomic fragments for the presence of genes or gene families; or for bacterial diagnostics by examining the ribosomal RNA genes; or for viral diagnostics by scanning for the presence of viral nucleic acid sequences in a sample. SPADT may also be used in forensic medicine by detecting and identifying species specific sequences or for the presence of major histocompatibility complex. The present sequence represents a primer specific for the human harpesvirus 6 (HHV6) p41 gene. The primer is used in an example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antiinicrobial; gene therapy; vaccine; amylase; cancer; amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factor; complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection; nervous system disease; ss.
The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoietin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histones, kinases, colony stimulating factors, complement related proteins, cytochromes, kinesins, cytokines, interferons, interleukins, Gytochromes, kinesins, cytokines, interferons, interleukins, G-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded
                                                                                                                                                                                 Polymorphic nucleic acids encoding e.g. amylases, oncogenes and histones, useful for diagnosing and cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2568 ttcttcttctttttttttttctgaaaaaaggaaaaa
                                                                                                                                                   Claim 1;
                                                                                                                                                                                                                                                                                                                                                                       28-DEC-1999;
27-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 40 BP; 9 A; 2 C; 4 G;
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27; Conser
                                                                                                                                                                                                                                                                                                                                     CURAGEN CORP
                                                                                                                                                                                                                                                                                                 RΑ,
                                                                                                                                               Page 2369; 4143pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                         2000US-0173419.
                                                                                                                                                                                                                                                                                                 Leach M;
                                                                                                                                                                                                                                                                                                                                                                                            99US-0173419
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Pred. No. 2.6e+04;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                    cyclins, polymerases, treating, e.g.
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AAQ87804
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Best Local S
Matches 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, leukaemia), diseases of the nervous system and an infection of pathogenic
                                   used to detect genes associated with allergic disease (diseases involving mutations in the beta chain gene). They may be used in neonatal screening, prenatal diagnosis etc. This sequence is an IgE ligand which is immobilised on a support and used to bind to probe/IgE receptor gene conjugates. The probe is then eluted by a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3473 ttgctattttttttttcataatttactatttatgatgtatttaa
                                                                                             DNA probes (See AAQ87799-803) having sequences identical or complementary to parts of the immunoglobulin E (IgE) receptor, are
                                                                                                                                                                                                           WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ87804 standard; DNA;
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                                                                                                                                Claim 8; Page 8; 17pp; English.
                                                                                                                                                                     Probes for mutation(s) in receptor - for the diagnos
                                                                                                                                                                                                                                                                                 22-OCT-1993;
                                                                                                                                                                                                                                                                                                                                 26-APR-1995
                                                                                                                                                                                                                                                                                                                                                        EP649910-A.
                                                                                                                                                                                                                                                                                                                                                                                                                    misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IgE receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-NOV-1995
                                                                                                                                                                                                                                                        (SUME ) SUMITOMO ELECTRIC IND CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    them may be used in the prevention, diagnosis and treatment of seases associated with inappropriate expression of the proteins listed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunoglobulin;
                                                                                                                                                                                                                                 Nakata M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              screening; diagnosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  beta chain ligand
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                                                                                                                                                                                                                                                                                 93JP-0265144.
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                                                                                                                                                                    diagnosis
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                                                                                                                                                                                                                                 Osoegawa M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IgE; receptor; beta; allergic disease;
diagnosis; ss.
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                                                                                                                                                                                                                                                                                                                                                                               "May optionally be labelled (product of ABI company)"
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                                                                                                                                                                                  beta chain
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                                                                                                                                                                      of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31 T; 0 other;
                                                                                                                                                                     allergic
                                                                                                                                                                                                                                 Ra
                                                                                                                                                                                  gene of a high affinity
                                                                                                                                                                                                                                 'n
                                                                                                                                                                      disease,
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Sequence

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BP;

12 A; 17

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probe/IgE receptor gene conjugates.
gradual rise in temperature.

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RESULT 1
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Best Local
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Best Local Similarity
                                                                                                                                                                                                                                                              A polypeptide having trypsin inhibitory activity, mol.wt. 22-28 k or 17-23 kD (as determined by SDS-PAGE under reducing or non-reducing conditions, respectively) and an N-terminal sequence as AAR79913 is claimed. The polypeptide was isolated from human T980 cells and is useful for treating pancreatitis, shock, multiple organ failure and disseminated intravascular coagulation. The present sequence is that of an antisense primer used for cloning
                                                                                                                                                 3467 gaattettgctatttttttttcataatttactatttatgatgtatt 3512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Trypsin; inhibitor; human T98G cells; pancreatitis; shock; DIC; multiple organ failure; disseminated intravascular coagulation; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2256 cagagagttgaggaccgcagcatcccctccactccagagttgg
Human SNP oligonucleotide #7280.
                        24-JAN-2002
                                                                    AAL34072 standard;
                                                                                                                                                                                                                               Sequence 48 BP; 2 A; 1 C; 1 G; 44 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                     Example 7; Page 15; 25pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-MAR-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               JP07242700-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Trypsin inhibitory protein cDNA antisense primer-1.
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                                                                                                                            pancreatitis,
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                                                                                                                                                                                                                                                      cDNA coding
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                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                           having trypsin inhibitory activity - for the treatment itls, shock, multi-organ failure, etc.
                                                                                                                                                                         Conservative
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                      (first entry)
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                                                                    DNA;
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                                                                    ВP
                                                                                                                                                                       Score 22; DB 16;
Pred. No. 3.3e+04;
0; Mismatches 15
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Pred. No. 3
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                                                                                                                                                                                              Length 48;
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AAL28913
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Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  variants of proteins related to amylases, amyloid proteins, angio
apoptosis related proteins, cadherin, cyclin, polymerase, oncogen-
histores, kinases, colony stimulating factors, complement related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; neuroprotective; antimicrobial; gene therapy; vaccine; amylase; cancer amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating fact complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation;
                                                                                                                                                                                                                        2564 agctttcttcttttttttttttctgaaaaaaggaaaaagcacaca 2609
Human SNP oligonucleotide #2121.
                                                                                                       AAL28913 standard;
                                                                                                                                                                                                                                                                                                                                                Sequence 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoietin,
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                                   24-JAN-2002
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                                                                                                                                                                                          l Similarity
31; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ctorial disease; autoimmune disease; infection; system disease; ss.
                                                                                                                                                                                                                                                                                                                                                BP; 17 A; 5 C; 7 G;
                                                                                                                                                                                                                                                              Conservative
                                  (first entry)
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                                                                                                       DNA;
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                                                                                                       ВP
                                                                                                                                                                                                                                                          Score 22; DB Pred. No. 3.4e 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                T; 0 other;
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Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic;

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AAH20339
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Best Local S
Matches 26
                                                                                                                                                                                                                                                                                                                                                                                                              proteins, cytochromes, kinesins, cytokines, interferons, interleukins, 6-protein coupled receptors and thioesterases. The present sequence is one such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of diseases associated with inappropriate expression of the proteins listed above. Disorders that may be prevented, diagnosed and/or treated include multifactorial diseases with a genetic component, such as autoimmune diseases (e.g. rheumatoid arthritis, multiple sclerosis, diabetes, systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                amyloid protein; angiopoletin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating factor; complement related protein; cytochrome; kinesin; cytokine; interferon; interleukin; G-protein coupled receptor; thioesterase; inflammation; multifactorial disease; autoimmune disease; infection; nervous system disease; ss.
                                                                                                                                                                                                                                   2007 ccttggaaaagcttgaagttgcagagtcagagt 2039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to oligonucleotides encoding polymorphic variants of proteins related to amylases, amyloid proteins, angiopoietin, apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, histores, kinases, colony stimulating factors, complement related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g.
                                                    HHV6 virus p41 gene specific primer p41FH46 SEQ ID
                                                                                     01-AUG-2001
                                                                                                                                                                                                                                                                                                                                                        Sequence 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 1988; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Shimkets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200147944-A2
                           Primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-DEC-1999; 99US-0173419
27-DEC-2000; 2000US-0173419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-DEC-2000; 2000WO-US35498
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                              AAH20339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CURA-) CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                                                  leukaemia), diseases
                                                                                                                                                                                                                       15
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                                                                                                                                                                                                                     ccttggaaaagccgcaagttgcttactccgagt 47
                                                                                                                                                                                                                                                                               Similarity
26; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        autoimmune diseases and infections
                          solid phase amplification of DNA template;
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                                                                                                                                             standard;
                                                                                                                                                                                                                                                                                                                                                        BP;
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                                                                                     (first
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                                                                                                                                                                                                                                                                                                                                                        13
            bacterial diagnostic;
                                                                                                                                             DNA;
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                                                                                   entry)
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Pred. No. 3
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            p41;
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            HHV6;
                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                   infection
            SPADT;
                                                       20
                                                                                                                                                                                                                                                                                                            50;
                          detection;
                                                                                                                                                                                                                                                                                                                                                                                              of pathogenic
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CC specific nucleic acid in a sample containing DNA. The method comprises CC princers are irreversible bound to a solid support, and the DNA from a CC primers are irreversible bound to a solid support, and the DNA from a CC sample is absorbed and reversibly bound, incubated under amplification CC reaction conditions and the presence of the specific target DNA is CC detected. The method is useful for detecting the presence of a specific CC nucleic acid (e.g. bacterial, viral or parasitic DNA) in a sample or in a CC cell. SPADT may be used for scanning large genomic fragments for the CC presence of genes or gene families; or for bacterial diagnostics by scanning the ribosomal RNA genes; or for bacterial diagnostics by scanning the presence of viral nucleic acid sequences in a sample. SPADT may CC also be used in forenesic medicine by detecting and identifying species specific sequences or for the presence of major histocompatibility complex. The present sequence represents a primer specific for the human illustrating the method of the invention.
                        Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting the presence of a specific nucleic acid in a sample containing DNA, useful in scanning large genomic fragments for presence of genes or gene families, comprises performing solid amplification of DNA template
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rovera G,
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                        Similarity
                                                                                                                        40
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                                                                                                                        BP;
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Score 21.6; DB 22
Pred. No. 3.9e+04;
0; Mismatches 9
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                                                                                                                        other;
                                              22;
                                            Length
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phase
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RESULT 14
AAL31293/c
                                                                                           Matches
AAL31293 standard; DNA; 50
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                                                1 ttttttttttttttttttttcgggaacatagagaaa
                                                           tttcttcttctttttttttttctgaaaaaaggaaaaa
                                                                                            27;
                                                                                           Conservative
ВP
                                                                                           0,:
                                                                      2602
                                                                                           9;
                                                                                           Indels
                                                                                           0;
                                                                                           Gaps
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B Q

multifactorial disease; autoimmune disease; information nervous system disease; ss. neuroprotective; antimicrobial; gene therapy; vaccine; amylase; can amyloid protein; angiopoietin; apoptosis related protein; cadherin; cyclin; polymerase; oncogene; histone; kinase; colony stimulating f complement related protein; cytochrome; kinesin; cytokine; interfer Human SNP Immunosuppressive; immunostimulatory; antiinflammatory; cytostatic; oligonucleotide #4501. (first entry)

cancer;

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RESULT 15
AAV83644/c
                                                                                                                                                                                                                                                                                                                                                           Qy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to oligonucleotides encoding polymorphic cc variants of proteins related to amylases, amyloid proteins, anglopoietin, cc apoptosis related proteins, cadherin, cyclin, polymerase, oncogenes, fistones, kinases, colony stimulating factors, complement related cc proteins, cytochromes, kinesins, cytokines, interferons, interleukins, cc g-protein coupled receptors and thioesterases. The present sequence is cone such oligonucleotide. The oligonucleotides and the peptides encoded by them may be used in the prevention, diagnosis and treatment of cc diseases associated with inappropriate expression of the proteins listed cabove. Disorders that may be prevented, diagnosed and/or treated include cc multifactorial diseases with a genetic component, such as autoimmune cc systemic lupus erythromatosus and Grave's disease), inflammation, cancer (e.g. cancers of the bladder, brain, breast, colon and kidney, cancer cc creanisms), diseases of the nervous system and an infection of pathogenic
XXX PF XXX
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                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 27; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polymorphic nucleic acids encoding e.g. amylases, cyclins, polymerases, oncogenes and histones, useful for diagnosing and treating, e.g. cancer, autoimmune diseases and infections -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-465210/50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-DEC-1999; 99US-0173419.
27-DEC-2000; 2000US-0173419.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                Claim 1; Page 2680; 4143pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Shimkets RA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 50 BP;
                                                                                                                             DNA
                                                                                                                                                                       Oligonucleotide
                                                                                                                                                                                                   01-MAR-1999
                                                                                                                                                                                                                                 AAV83644;
                                                                                                                                                                                                                                                            AAV83644 standard; DNA; 35 BP.
                                                                     W09845476-A1
                                                                                               Synthetic
                                                                                                                                         Repetitive
            08-APR-1998;
                                          15-OCT-1998.
                                                                                                                           etitive sequence; carcinogenic;
instability; cancer; diet; prin
                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                    (first entry)
               98WO-GB00869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24 A; 5 C; 4 G; 17 T; 0 other;
                                                                                                                                                                       used in the construction of assay plasmids.
                                                                                                                                                                                                                                                                                                                                                                                                         0.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                       Score 21.6;
Pred. No. 4.
                                                                                                                               primer;
                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                              human dietary component;
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                                                                                                                                                                                                                                                                                                                                                                                                          4.4e+04;
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                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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The present sequence represents an oligonucleotide used in the CC construction of assay plasmids, which are used in the course of the CC invention. The specification describes an assay for testing the CC carcinogenic properties of a test substance. The assay comprises CC introducing into cells a reporter gene expression vector comprising a CC repetitive DNA sequence which exhibits instability in cancer cells, CC whereby instability of the repetitive DNA sequence affects expression of CC the reporter gene, exposing the resulting cells to the test substance CC and determining whether the test substance is carcinogenic or CC anti-carcinogenic by comparing the frequency of reporter gene expression in CC in the resulting cells with the frequency of reporter gene expression in CC cells which have not been exposed to the test substance. The assay can CC cells which have not been exposed to the test substance. The assay can CC cells which have not been exposed to the test substance of instability, and therefore some types of cancer, and can be used to CC contribute to the scientific basis for a healthy diet.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Assay for testing the carcinogenic properties of a test substance by introduction of a reporter gene expression vector containing a repetitive DNA sequence that is unstable in cancer cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-024011/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 17; 103pp; English.
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Sequence
  35
  BP;
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     22 A; 5 C; 6 G; 2 T; 0 other;
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Search completed: September 18, Job time: 8329 sec 2002, 01:22:27 밁 Ş

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2561 cccagctttcttcttcttttttttttttctgaa 2591

CCGGGCTTTTTTTTTTTTTTTTTTTTTCCGAA 3

Matches Query Match Best Local 9

25;

Conservative

0; Pred. Score 21.4; Mismatches No.

Similarity

0.6%;

4e+04; DВ 20; 6;

Length 35; Indels

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Gaps

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Result
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Listing first 45 summaries
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Perfect score:
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1: /cgn2_6/ptodata/1,
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5: //gn2_6/ptodata/1,
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3713
      GenCore version
Copyright (c) 1993 - 2000
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/BCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/backfiles1.seq:*
        Length
    ВG
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US-08-718-904-50
PCT-US95-10973A-43
US-09-306-290-21
US-09-306-290-20
US-08-686-599A-14
US-08-52-623-14
US-08-52-623-14
PCT-US93-03256-14
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PCT-US93-03256-17
US-08-208-486-57
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US-08-208-486-57
US-08-171-389-456
US-08-171-389-476
US-08-171-389-577
US-08-171-389-577
US-08-173-389-576
US-08-173-3936-456
US-08-475-228A-456
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           Sequence 50, 7
Sequence 21, 7
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Sequence 17, 7
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Sequence 17, 7
Sequence 17, 7
Sequence 57, 8
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Sequence
50, Appl
21, Appl
20, Appl
14, Appl
14, Appl
35, Appl
35, Appl
37, Appl
17, Appl
18, Appl
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Query Match Best Local S Matches 28	US-08-718-904-50  US-08-718-904-50  Sequence 50, Applicat	2 2 2	28 20.2 29 20.2 30 2.2 31 20 32 20 33 20 34 20 35 20 36 20 37 20 38 20 41 20 41 20
h Similarity 75. 28; Conservative	G PATHON BEOND OF RESEARCH ON PARK S.		00000000000000000000000000000000000000
.6%; Score 22.6; .7%; Pred. No. 1.7 e 0; Mismatches	701 700 700 700 700 700 700 700 700 700	1 US 1 US	1 1 1 1 1 2 3 3 8 8 4 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
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Length 45; Indels 0; (	ACIDS AND	Sequence Sequence Sequence	
Gaps 0;	LIGANDS FOR	5, Appli 5, Appli 5, Appli	576, App 577, App 264, Appl 22, Appl 21, Appl 5, Appl 5, Appli 5, Appli 5, Appli 5, Appli

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                                                                                                                                                                                                                       US-09-306-290-21
                                                                                                                                                                                                                                            RESULT
                                                                                                                                                                   Sequence 21, Application US/09306290 Patent No. 6221635 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.6%;
Best Local Similarity 75.7%;
Matches 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 45 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 43, Application PC/TUS9510973A GENERAL INFORMATION:
             APPLICANT: Rovera, Giovanni
APPLICANT: Mukhopadhyay, Sunil
APPLICANT: Mukhopadhyay, Sunil
TITLE OF INVENTION: METHODS FOR SOLID-PHASE AMPLIFICATION OF DNA TEMPLATE
TITLE OF INVENTION: (SPADT) USING MULTIARRAYS
FILE REFERENCE: 09924-10
CURRENT FILING DATE: 1999-05-06
CURRENT FILING DATE: 1999-05-06
NUMBER OF SEQ ID NOS: 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Nottenburg, Carol
REGISTRATION NUMBER: 39,317
REFERENCE/DOCKET NUMBER: 760.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Prizm Pharmaceuticals, Inc.
TITLE OF INVENTION: CONJUGATES OF VASCULAR ENDOTHELIAL GROWTH FACTOR WITH TARGET NUMBER OF SEQUENCES: 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: DNA (genomic)
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LOCATION:
NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Seattle
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                                                                                                                                                                                                                                                                                               3 ATGGCCTCGTCGTCGTCGGCCTCGTCGTCGGCCG 39
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nucleic acid
DEDNESS: single
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6300 Columbia Center, 701 Fifth Avenue
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Pred. No. 1.
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; OTHER INFORMATION: Description of Artificial Sequence: Primer p41; OTHER INFORMATION: FH46
US-09-306-290-20
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; OTHER INFORMATION:
US-09-306-290-21
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CURRENT FILING DATE: 1999-05-06
NUMBER OF SEQ ID NOS: 43.
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 21
LENGTH: 40
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Best Local S
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Best Local Similarity
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APPLICANT: Mukhopadhyay, Sunil
TITLE OF INVENTION: METHODS FOR SOLID-PHASE AMPLIFICATION OF DNA TEMPLATE
TITLE OF INVENTION: (SPADT) USING MULTIARRAYS
FILE REFERENCE: 09924-10
                                                                                                                                                                                                                                          GENERAL INFORMATION: APPLICANT: Olsson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
COUNTRY: USA
ZIP: 22314-3187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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APPLICANT:
                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                            APPLICANT: Donelson, John E.
TITLE OF INVENTION: LYMPHOCYTE STIMULATING FACTOR
                                                                                                                                                                                APPLICANT:
                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
nes 27; Conserv
                                                                STATE:
                                                                                   CITY:
                                                                                              STREET:
                                                                                                               ADDRESSEE:
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                                                                               Alexandria
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                                                                                              8: BURNS, DOANE, SWECKER & 699 Prince Street
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Bakhiet, Abdel-Moiz
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77.1%;
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Pred. No. 3
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 14, APF-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.6%;
Best Local Similarity 73.0%;
Matches 27; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Duffett, Benton S.
REGISTRATION NUMBER: 22,030
REFERENCE/DOCKET NUMBER: 00:
INFORMATION FOR SEQ ID NO: 14:
                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2567 tttcttctttttttttttctgaaaaaggaaaaag 2603
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ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
          NAME: Brainard, Thomas D.
REGISTRATION NUMBER: 32,459
REFERENCE/DOCKET NUMBER: 511
TELECOMMUNICATION INFORMATION:
                                                                              APPLICATION NUMBER: US/07/863,553
FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: |
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Abbott Laboratories
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: METHOD AND DEVICE FOR DETECTION OF TITLE OF INVENTION: NUCLEIC ACID OR ANALYTE USING TOTAL INTERNAL REFLECTION NUMBER OF SEQUENCES: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                            CLASSIFICATION: 435
                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                         STREET: One Abbott Park Road CITY: Abbott Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          42 TTTTTTTTTTTTTTTTAGTGAAATGACAATAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: TLTF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/0 FILING DATE: 26-JUL-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                       60064-3500
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EDNESS: unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bouma, Stanley R. Pabich, Edward K.
                                                                                                                                                                                                                                                                                                                                                                      United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CDS
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(708) 937-4884
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                                5158.US.01
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PCT-US93-03256-14
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Best Local
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 32 base pairs
TYPE: nucleic acid
                                                                                                                                                           TELEFAX: (708) 937-26 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 51
TELECOMMUNICATION INFORMATION:
TELEPHONE: (708) 937-4884
                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/0
FILING DATE: 06 APRIL 19
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                            MOLECULE TYPE: DNA (genomic) FEATURE:
                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn and WordPerfect
CURRENT APPLICATION DATA:
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                                                                                                                                            SEQUENCE CHARACTERISTICS:
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              NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: Unc. Park CITY: Abbott Park
                                                                             STRANDEDNESS:
TOPOLOGY: li
                                                                                                                                                                                                                                       REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                      FILING DATE: 1
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
TOPOLOGY: lir
                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE:
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                                                                                                                          LENGTH:
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25; Conserv
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                                                                                                        NUCLEIC ACID
                                                                                                                                                                                                                                                        Brainard, Thomas D.
                                                                                                                        32 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              One Abbott Park Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Abbott Laboratories
VENTION: METHOD AND DEVICE FOR DETECTION OF
VENTION: NUCLEIC ACID OR ANALYTE USING TOTAL INTERNAL REFLECTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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              misc_feature
32
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32
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Pred. No. 4.7e+03
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Query Match
Best Local Similarity 78.3
Matches 25; Conservative

0.6%;

Score 20.8; DB 5; Pred. No. 4.7e+03; 0; Mismatches 7;

Length 32;

Indels

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Gaps

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-389-341-65/c
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US-08-316-439A-35
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TITLE OF INVENTION: ESSENTIAL GENES OF YEAST AS TARGETS FOR ANTIFUNGAL
TITLE OF INVENTION: AGENTS, HERBICIDES, INSECTICIDES AND ANTI-PROLIFERA
TITLE OF INVENTION: DRUGS
FILE REFERENCE: 9301-057
CURRENT APPLICATION NUMBER: US/09/389,341
CURRENT FILING DATE: 1999-09-02
EARLIER APPLICATION NUMBER: 09/315,794
EARLIER FILING DATE: 1999-05-21
NUMBER OF SEQ ID NOS: 72
SOFTWARE: PATENTIN VET: 2.0
SEQ ID NO 65
LENGTH: 45
THERE FILING ASSET IN THE PROPERTY OF THE P
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Best Local S
Matches 29
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Patent No. 6200803
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 35, Application US/08316439A Patent No. 5840520 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Artificial Sequence
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MEDIIM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: CLARKE, DAVID KIRKWOOD APPLICANT: PALESE, PETER M TITLE OF INVENTION: RECOMBINANT NITITLE OF INVENTION: SYSTEMS
                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/190,678
FILING DATE: February 1, 1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
nes 29; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: PALO ALTO
STATE: CALIFORNIA
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                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/316,439A FILING DATE: September 30, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET:
   FILING DATE:
APPLICATION NUMBER: 07/925,0
FILING DATE: August 4, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94306
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: COOLEY GODWARD CASTRO HUDDLESON & TATUM FIVE PALO ALTO SQUARE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
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                                        07/925,061
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. 6.8e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 45;
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FILING DATE: August 28, 1989
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: CSER, LUANN
REGISTRATION NUMBER: AVIR-010/00US
REFERENCE/DOCKET NUMBER: AVIR-010/00US
TELEPHONE: (415) 843-5165
TELEPHONE: (415) 843-5165
TELEPAX: 380816 COLLEY PA
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: US-08-316-439A-35
                                             US-08-208-486-57/c; Sequence 57, Application US/08208486; Patent No. 5389531; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   В
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                                                                                                          RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: FOSTER, DONALD C.; MULVIHILL, EILEEN R.; O'HARA, ; PATRICK J.; PINGEL, KURT; YOSHITAKE, SHINJI TITLE OF INVENTION: THROMBIN-ACTIVATED TISSUE PLASMINOGEN
                                                                                                                                                                                                                                                                                                                          FILING DATE: 22-MAY-1987: SEQ ID NO:28:
                                                                                                                                                                                                                                                                                                                                                                                                           ACTIVATORS
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Best Local Similarity
Matches 30; Conserv
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Best Local
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: No. 584
                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 34
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICANT: Ito, Junetsu
APPLICANT: Yoo, Seung-Ku
TITLE OF INVENTION: METHODS TO REPLICATE DNA in vitro USING
                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
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25; Conser
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DEDNESS: single
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No. 5840520ember 21,
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                                                                                                                                                                                                                     Score 20.2; D
Pred. No. 7.4e
0; Mismatches
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                                                                                                                                                                                                                                   .4e+03;
                                                                                                                                                                                                                                                    DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 2;
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; MOLECULE TY:
US-08-208-486-57
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Matches
                                                                                                                                                                                                                                                                                                                                                     Sequence 17, Application US/08284784 Patent No. 5629172
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                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (602) 495-9475
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: BOULD COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 1.2 Mb
COMPUTER: Packard Bell (IBM PC/AT compatible)
COMPUTER: MS-Dos, Version 5.0
                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA: 07/869,916
APPLICATION NUMBER: 07/869,916
FILING DATE: APPLICATION NUMBER: Japan 2405;
FILING DATE: August 26, 1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: MS-Dos, Version SOFTWARE: WordPerfect Version 5.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Cahill, Sui
STREET: 155 Park One,
                                                                                                                                                                                    TITLE OF INVENTION: EXPRESSION OF FUSION POLYPEPTIDES
TITLE OF INVENTION: TRANSPORTED OUT OF THE CYTOPLASM WITHOUT LEADER SEQUENCES: 44
                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                           APPLICANT: OLSEN, DAVID R. APPLICANT: CARRILLO, PEDRO A.
                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            122 gcttactcttctgtgaaagggaaagtatcccc 154
                                                                                                                                                                                                                                                                                                                                                                                                                                                        33 GCTTAGTGTTTTGGGGGAGGGGCACGTATCCCC 1
                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 75.8 nes 25; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS:
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REFERENCE/DOCKET NUMBER: #3954-A-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Janelle Faunce Raupp REGISTRATION NUMBER: 30,48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 435
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   OPERATING SYSTEM:
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                                                                                                                                                         ADDRESSEE:
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                                                                                                                        Palo Alto
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TYPE: Other nucleic acid (synthetic DNA)
                                                                                                      California
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                                                                                                                                         E: MORRISON & FOERSTER
755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       40 base pairs
                                                                                                                                                                                                                                                             OLSON, PAMELA S. OLSEN, DAVID R.
                                                                                     USA
                                                                                                                                                                                                                                                                                                ZHANG, YANG
                                                                                                                                                                                                                                                                                                                  MASCARENHAS, DESMOND
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PC-DOS/MS-DOS
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Pred. No. 8
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Patent No. 5914254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.5%;
Best Local Similarity 68.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (415) 494-079
TELEX: 706141
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1232 cctgatggtcttttttttggatgctgacatgtacaacgctct 1272
                                                                        APPLICATION NUMBER: 08/284,784
FILING DATE: 02-AUG-1994
APPLICATION NUMBER: 08/100,744
FILING DATE: 02-AUG-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                            CURRENT APPLICATION DATA: APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 35,636
REFERENCE/DOCKET NUMBER: 35,636
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813,5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: PARK, FREDDIE K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                         PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM
                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 46 base pairs
TELECOMMUNICATION INFORMATION: TELEPHONE: 650-813-5600
                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 02-AUG-1994
CLASSIFICATION: 530
                             NAME: Buffinger, Nicholas S
REGISTRATION NUMBER: 39,124
REFERENCE/DOCKET NUMBER: 220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: (415) 494-0792
                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                               SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                                                            OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                 COMPUTER:
                                                                                                                                                                                                                                                                                               MEDIUM TYPE:
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STREET: 755 PAGE MILL ROAD
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94304-1018
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                                                                                                                                                                                                                                                                              IBM Compatible
                                                                                                                                                                                         12-MAY-1997
                                                                                                                                                                                                                                                                                               Diskette
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TRANSPORTED OUT OF THE CYTOPLASM WITHOUT LEADER
                                                                                                                                                                                                                                                                  Windows
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                                22095-20275.21
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Pred. No. 9.1e+03;
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US-08-171-389-456/c
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Best Local Similarity
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                                   TELEFAX: (415) 324-0960 INFORMATION FOR SEQ ID NO: 4 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1232 cctgatggtcttttttttggatgctgacatgtacaacgctct 1272
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LENGTH: 46 base pairs
                                                                                                           REGISTRATION NUMBER: 33,875
REFERENCE/DOCKET NUMBER: 4600-0175/G19P3
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: US 08/123,936
FILING DATE: 17-SEP-1993
PRIOR APPLICATION NUMBER: US 07/996,783
PRIOR APPLICATION NUMBER: US 07/996,783
FILING DATE: 23-DEC-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                              APPLICATION NUMBER: US 07/723,618 FILING DATE: 27-JUN-1991 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Sequence-Directed DNA Binding TITLE OF INVENTION: Molecules, Compositions and M NUMBER OF SEQUENCES: 641
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APPLICANT: Cantor, Charles R.
                                                                                                                                                                                       FILING DATE: 22-JUN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
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APPLICANT: Fry, Kirk E.
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APPLICANT: Turin, Lisa M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Redwood City
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/081,070
                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/171,389
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                                                                                               TELEPHONE:
                       LENGTH:
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nucleic acid
                                                                                                                                                                       Fabian, Gary R.
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                       50 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Genelabs Technologies, Inc.
                                                                                               (415) 324-0880
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Pred. No. 9.1e+03;
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; INDIVIDUAL ISOLATE: Human beta-LH gene US-08-171-389-456
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Matches
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                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/
EFILING DATE: 27-JUN-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/
EFILING DATE: 22-JUN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                              TELEFAX: (415) 324-09 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORIGINAL SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/123,936
FILING DATE: 17-SEP-1993
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TITLE OF INVENTION: Sequence-Directed DNA Binding
TITLE OF INVENTION: Molecules, Compositions and Methods
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                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 17-SEP-1993
PRIOR APPLICATION DATA:
, APPLICATION NUMBER: US 0'
FILING DATE: 23-DEC-1992
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APPLICATION NUMBER: US
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                         TOPOLOGY: li
                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 324-0880
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 63.3 nes 31; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Genelabs Technologies, Inc. STREET: 505 Penobscot Drive
                                                                                                                                                           TELEPHONE:
                                                                                                                                                                                              REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                            NAME: Fabian, Gary R. REGISTRATION NUMBER:
                                                            STRANDEDNESS: double
                                                                                            LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               576, Application US/08171389
o. 5578444
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                                                                            nucleic acid
                                                                                              50 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Andrews,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PatentIn Release #1.0, Version #1.25
                                               linear
                NO
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                              DNA (genomic)
                                                                                                                                             324-0960
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                                                                                                                                                                                                              33,875
                                                                                                                                576:
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                                                                                                                                                                                              4600-0175/G19P3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 20.2; DB 1;
Pred. No. 9.6e+03;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
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Sequence:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Post-processing: Minimum Match
Maximum Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OM nucleic - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Database
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 1.0
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3713
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pending_Patents_NA_Main: *
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/cgn2_6/ptodata/1/pna/USO16_COMB.seq:*
/cgn2_6/ptodata/1/pna/USO17_COMB.seq:*
/cgn2_6/ptodata/1/pna/USO18_COMB.seq:*
/cgn2_6/ptodata/1/pna/USO181_COMB.seq:*
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/cgn2_6/ptodata/1/pna/USO185_COMB.seq:*
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/cgn2_6/ptodata/1/pna/US092_COMB.seq:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /cgn2_6/ptodata/1/pna/US086_COMB.seq:*
/cgn2_6/ptodata/1/pna/US087_COMB.seq:*
/cgn2_6/ptodata/1/pna/US088_COMB.seq:*
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Compugen Ltd
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44: /cgn2_6/ptodata/1/pna/US6005_COMB.seq: *
45: /cgn2_6/ptodata/1/pna/US6006_COMB.seq: *
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63: /cgn2_6/ptodata/1/pna/US6023_COMB.seq: *
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67: /cgn2_6/ptodata/1/pna/US6029_COMB.seq: *
68: /cgn2_6/ptodata/1/pna/US6032_COMB.seq: *
70: /cgn2_6/ptodata/1/pna/US6032_COMB.seq: *
71: /cgn2_6/ptodata/1/pna/US6032_COMB.seq: *
72: /cgn2_6/ptodata/1/pna/US6032_COMB.seq: *
73: /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
74: /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
75: /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
76: /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
76: /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
77: /cgn2_6/ptodata/1/pna/US6033_COMB.seq: *
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Result No.

Score

Query Match Length DB

ID

Description

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27 28 29 30 31	223 24 26	13 14 15 16 17 17 18 19 20	1 2 3 3 4 4 4 7 7 7 9 9 110 110 121
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12 18 23 33	17666	29 7 8 8 8 14 14 64	37 29 17 17 29 29 29 29
US-08-805-382-29 US-08-805-383-29 US-09-449-249-50 US-09-617-325-43 US-09-861-257-29	US-08-213-446A-43 US-08-213-447A-43 US-08-297-961A-43 US-08-305-771A-43 US-08-305-771A-43	US-09-755-374A-13433 US-08-346-731-6670 US-08-420-856-6670 US-08-420-856A-6670 US-08-420-856A-6670 US-09-78-617-2126 US-09-783-590-6670 US-09-783-590-6670 US-09-783-523-1976	US-10-003-354-6 US-09-755-374A-25691 US-09-755-374A-1769 US-09-342-217-44074 US-09-755-374A-1769 US-09-396-196F-83096 US-08-401-791A-15261 US-08-401-791A-15261 US-08-401-791A-15261 US-08-401-791A-15261 US-08-401-791A-15261 US-09-755-374A-2381 US-09-755-374A-2381 US-09-755-374A-6837
Sequence 29, Appl Sequence 29, Appl Sequence 50, Appl Sequence 43, Appl Sequence 29, Appl	Appl Appl Appl Appl Appl	13433, 5670, A 5670, A 5670, A 6670, A 2126, 6670, 1976,	Sequence 6, Appli Sequence 25691, A Sequence 4074, Ap Sequence 17699, A Sequence 17699, A Sequence 15261, A Sequence 15261, A Sequence 2381, Ap Sequence 25692, A Sequence 5024, Ap Sequence 5024, Ap

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US-10-003-354-6
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                                                                                                                                                                                                                                                                                                                                                                                                                             ; FEATURE:
; OTHER INFORMATION: PCR Probe
US-10-003-354-6
                                                                                                                                                                                                                                                                    US-09-755-374A-25691
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Best Local S
Matches 26
                                                                                                  PRIOR APPLICATION NUMBER: 60/174962
PRIOR FILING DATE: 2000-01-07
NUMBER OF SEQ ID NOS: 28742
SEQ ID NO 25691
LENCTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 6, Application US/10003354
GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: Susan M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION
TITLE OF INVENTION: EXPRESSION
FILE REFERENCE: RTS-0348
                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 6
LENGTH: 26
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                            Sequence 25691, Application US/09755374A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/003,354
CURRENT FILING DATE: 2001-12-06
NUMBER OF SEQ ID NOS: 89
                                                                                                                                                                                APPLICANT: Leach, Martin
APPLICANT: Shimkets, Richard A.
TITLE OF INVENTION: Nucleic Acids Containing
TITLE OF INVENTION: Use Thereof
FILE REFERENCE: 15966-611
                                                                                                                                                        CURRENT APPLICATION NUMBER: US/09/755,374A
CURRENT FILING DATE: 2001-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature
LOCATION: (25)...(0)
COTHER INFORMATION: 1 of 2 allelic variants (25692 is other entry)
                                                                       TYPE: DNA
ORGANISM: Homo sapiens
        NAME/KEY: misc_feature
                                                             FEATURE:
LOCATION:
                                                                                              ENGTH: 50
                                                                                                                                                                                                                                                                                                                 0.7%;
Similarity 100.0%;
26; Conservative
(0)...(0)
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74
37
14
64
22
37
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US-09-726-173A-102
US-09-396-196F-83095
US-09-522-303-489
US-60-353-790-1521
US-10-017-974-23096
US-09-078-617-2930
US-60-253-378-27503
US-60-253-378-27503
US-09-572-021-4910
US-10-017-974-35315
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                                                                                                                                                                                                                                                                                                                                                                   0;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-798-074-3200
US-08-798-074B-3200
US-09-755-374A-2636
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                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OF
                                                                                                                                                                                                                                                                                                                                                                   DB 37; I
. 5.4e+04;
ches 0;
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                                                                                                                                                                                                           Single Nucleotide Polymorphisms and
                                                                                                                                                                                                                                                                                                                                                                                          Length 26;
                                                                                                                                                                                                                                                                                                                                                                    Indels
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Sequence 3200,
Sequence 2636,
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Sequence
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Sequence
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1521, Ap
23096, A
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35315,
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27503, A
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5, A
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                                                                         -US-09-342-217A-4074; Sequence 4074, Application US/09342217A
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                                                                                                                                                            QΥ
                                                                                                                                                                                                                                                US-09-342-217-4074
                                                             GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                Query Match
Best Local S
                                                                                                                                                                                      Matches
  TITLE OF INVENTION: Human Genes, Sequence FILE REFERENCE: PO-40 CURRENT APPLICATION NUMBER: US/09/342,217A CURRENT FILING DATE: 1999-06-29
                                                                                                                                                                                                                                                                                                                     TELEFAX: (301) 309-85
INFORMATION FOR SEQ ID NO:
                                                    APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: HP Vectra 48
OPERATING SYSTEM: MSDO
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Craig A. Rosen, TITLE OF INVENTION: Human
                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 50 base pairs.
                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                  2562 ccagctttcttcttcttttttttttttttgaaaaaaggaa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: 9410
CTTY: Rockville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                      13 CGAGTTTTTTTTTTTTTTTTTTTTNANAAAAANNAA 50
                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                               NAME: Benson, Bob
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                             TOPOLOGY: 1i
                                                                                                                                                                                                                                                                                    TYPE:
                                                                                                                                                                                                                                                                                                                                            TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                      . Similarity 73.7
28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20850
                                                                                                                                                                                                                                                                                 nucleic acid
                                                    Rosen, Craig, et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9410 Key
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                                                                                                                                                                                                                                                             linear
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                                        Human Genes, Sequences,
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5660
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                                                                                                                                                                                      Score 24.4; D
Pred. No. 2e+0
0; Mismatches
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Pred. No.
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3479 tttttttttcataatttactattatgatgtatttaagtgttttattaa 3527
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and Expression Products 40
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US-09-755-374A-17699; Sequence 17699, Application US/09755374A; GENERAL INFORMATION:
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; OTHER INFORMATION: n equals a,t,g, or
US-09-342-217A-4074
                                 ð
   밁
                                                                                                                                                            ; LOCATION: (0)...(0)
; OTHER INFORMATION: Accession number cg43932310
US-09-755-374A-17699
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Matches 28; Conserv
                                                                     Matches
                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: 60/174962
PRIOR FILING DATE: 2000-01-07
NUMBER OF SEQ ID NOS: 28742
SEQ ID NO 17699
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS:
SOFTWARE: Patentin V
SEQ ID NO 4074
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Leach, Martin
APPLICANT: Shimkets, Richard A.
APPLICANT: Shimkets, Richard A.
TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide Polymorphisms
TITLE OF INVENTION: Use Thereof
FILE REFERENCE: 15966-611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/755,374A CURRENT FILING DATE: 2001-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: 60/0 PRIOR FILING DATE: 1998-06-30
                                                                                                                                                                                                         NAME/KEY: misc_feature
LOCATION: (25)...(26)
OTHER INFORMATION: Nucleotide deleted between bases
NAME/KEY: misc_feature
LOCATION: (0) (0)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc feature LOCATION: (47)
OTHER INFORMATION: n equals a,t,g,
                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
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OTHER INFORMATION: n equals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                     LOCATION: (26)...(0)
OTHER INFORMATION: 1 of 2 allelic variants (17700 is
                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: n equals a,t,g,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc feature
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                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 50
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                  3470 ttcttgctattttttttttcataatttactatttatgatgtatt 3512
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ttttttttttttttttagcacttagtatttagcatttatt 49
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                                                                       Conservative
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73.7%;
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                                                                                      Score 23.8;
Pred. No. 2.
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Sequence 83096, Application Us
GENERAL INFORMATION:
APPLICANT: Michael Mittmann
APPLICANT: David Mack
APPLICANT: David Lockhart
APPLICANT: Affymetrix, Inc.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: 60/100,678
PRIOR FILING DATE: 1998-09-17
NUMBER OF SEO ID NOS: 127806
SOFTWARE: FastSEQ for Windows Version
SEO ID NO 83096
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-396-196F-83096
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GENERAL INFORMATION:
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Best Local (
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TITLE OF INVENTION: Methods of Genetic Analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: 3101.
              TELEFAX: (201) 994-1744 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 25
                                                                                                                                                                                                                                                                                                                                                                                      STREET: U.STREET: Noseland
CITY: Roseland
CTATE: New Jersey
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                                                                                                                  FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Olstein, Elliot
                                                                                                                                                                     CLASSIFICATION: 435
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                               COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS VES
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
                             REFERENCE/DOCKET NUMBER: 32:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (201) 994-1700
TELEFAX: (201) 994-1744
                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM: MEDIUM, TYPE: Diskette,
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF
                                                                                                                                                                                                                            APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U
                                                                                                    NAME: Olstein, Elliot REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                              6 Becker Farm Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Dillon, Patrick
Li, Haodong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Rosen, Craig
Ruben, Steve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Haseltine, William
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Carella, Byrne, Bain, Gilfillan, Cecchi, Stewart, & Olstein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human Genes, Sequences, and Expression Products
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.6%;
96.0%;
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                                                                                                                                                                                                                                               US/08/401,791A
                                                                                                    24,025
                                                                                                                                                                                                                                                                                                                                  3.50 inch, 1.4Mb
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                                                                                     325800-307
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                                                                                                                                                                                                                                                                                                                                    storage
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; TYPE: nucleic acid; STRANDEDNESS: double; TOPOLOGY: linear US-08-401-791A-15261
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RESULT 9
US-09-755-374A-2381
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US-08-401-791B-15261
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Best Local Similarity
                                                                                                                                             Query Match
Best Local Similarity
Matches 27; Conserv
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                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/401.
FILING DATE: MATCH 10, 1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: MICHELE M. Wales
REGISTRATION NUMBER: P-43,975
REGISTRATION NUMBER: P0-11
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8439
TELEPHONE: (301) 309-8439
                                                                                                                                                                                                                                                                                                   TELEFAX: (301) 309-843:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 50 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIIM TYPE: Diskette, 3.50 inch, 1.4Mb storage
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 6.2
SOFTWARE: ASCII Text
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2567 tttcttctttttttttttctgaaaaaaggaaaaa 2602
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TITLE OF INVENTION: Human Genes, Sequences and Expression Products -
NUMBER OF SEQUENCES: _17623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
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CITY: Rockville
STATE: Maryland
                                                                                                                                                                                                                                                              LENGTH: 50 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                         15 TTTTTTTTTTTTTTTTTTTNAAAAAAAAAAAAAA 50
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ilarity 75.0%;
Conservative
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Dillon, Patrick J.
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                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Li, Haodong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Haseltine, William A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                  linear
                                                                                                                                                           0.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Genome Sciences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US/08/401,791B
                                                                                                                                                                                                                                                                                                                                                                                                                            P-43,975
                                                                                                                                                                                                                                                                                                                                        15261:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 23.4; DB 8; Pred. No. 3.6e+05;
                                                                                                                                                           Score 23.4; DB 8;
Pred. No. 3.6e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9;
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                                                                                                                                                                            Length 50
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FEATURE:

NAME/KEY: misc_feature
LOCATION: (25)...(0)
OTHER INFORMATION: 2 of 2 allelic variants (25691 is other entry)
NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: Accession number cg44004721
US-09-755-374A-25692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
LOCATION: (25)...(26)
OTHER INFORMATION: Nucleotide deleted between bases
NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: Accession number cg43928213
US-09-755-374A-2381
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CURRENT APPLICATION NUMBER: 099/755,374A
CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: 60/174962
PRIOR FILING DATE: 2000-01-07
NUMBER OF SEQ ID NOS: 28742
SEQ ID NO 2381
LENGTH: 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 33; Conser
                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 28742
SEQ ID NO 25692
LENGTH: 50
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 25692, Application US/09755374A GENERAL INFORMATION:
APPLICANT: Leach, Martin APPLICANT: Shimkets, Richard A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2381, Application GENERAL INFORMATION:
                                                                                  Matches
                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Leach, Martin
APPLICANT: Shimkets, Richard A.
APPLICANT: Shimkets, Richard A.
TITLE OF INVENTION: Nuclecic Acids Containing Single Nucleotide Polymorphisms and
TITLE OF INVENTION: Use Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/755,374A
CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: 60/174962
PRIOR FILING DATE: 2000-01-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Nucleic Acids Containing TITLE OF INVENTION: Use Thereof FILE REFERENCE: 15966-611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: misc_feature
LOCATION: (26)...(0)
OTHER INFORMATION: 1 of 2 allelic variants (2382 is other entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2565 gctttcttcttcttttttttttctgaaaaaaggaaaaagcacacagcac 2613
                         3479 ttttttttcataatttactatttatgatgtatttaagtgttttattaa 3527
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                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.6%;
                                                                                                   0.6%;
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                                                                                  0;
                                                                                  Score 23.4; DB 29;
Pred. No. 3.6e+05;
0; Mismatches 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 23.4; DB 29; Pred. No. 3.6e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Single Nucleotide Polymorphisms and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16;
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                                                                                                                      Length 50;
                                                                                    Indels
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                                                                                  Gaps
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PCT-US01-47856-5024/c

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US-09-755-374A-6837/c

: Sequence 6837, Application US/09755374A

; GENERAL INFORMATION:
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APPLICANT: Altman, Petter
TITLE OF INVENTION: LEUKOCYTE EXPRESSION PROFILING
FILE REFERENCE: 506612000140
CURRENT APPLICATION NUMBER: PCT/US01/47856
CURRENT FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/241,994
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2000-10-6-08
PRIOR FILING DATE: 2001-06-08
NUMBER: US 60/296,764
NUMBER: DATE: 2001-06-08
NUMBER: DATE: 2001-06-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ORGANISM: Homo sapiens PCT-US01-47856-5024
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SEQ ID NO 5024
LENGTH: 50
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Best Local Similarity
                                                                                                                                                                                                                                                            SEQ ID NO 6837
LENGTH: 50
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                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 15966-611
CURRENT APPLICATION NUMBER: US/09/755,374A
CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: 60/174962
PRIOR FILING DATE: 2000-01-07
NUMBER OF SEQ ID NOS: 28742
                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Leach, Martin
APPLICANT: Shimkets, Richard A.
TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide Polymorphisms and Meth
TITLE OF INVENTION: Use Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: BIOCARDIA,
NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: Accession number cg44911096
                                                                     NAME/KEY: misc_feature
LOCATION: (25)...(26)
OTHER INFORMATION: Nucleotide deleted between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
                                                                                                                                                            NAME/KEY: misc_feature LOCATION: (26)...(0)
                                                                                                                                                                                                    FEATURE:
                                                                                                                                         OTHER INFORMATION: 1 of 2 allelic variants (6838 is
                                                                                                                                                                                                                         ORGANISM: Homo sapiens
                                                                                                                                                                                                                                               TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1214 gacttcttacaagacatccctgatggtcttttttttggatgctga 1257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              44 GAATGCCTATAAGAAATCTCAAAAGGTATTTGTTTTGGGTGCAGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Prentice, James
Phillips, Julie
Woodward, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Johnson,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wohlgemuth, Jay
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Quertermous, Thomas
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 23.2; DB Pred. No. 4e+05; O; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 50;
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                                                                            bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                         other entry)
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                                                                                                                                                                                                                                                                                                                                                             US-08-346-731-6670
                                                                                                                                                                                                                                                                                                                           Sequence 6670, Application US/08346731 GENERAL INFORMATION:
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Best Local Similarity
Matches 28; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 13433
LENGTH: 50
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local
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APPLICANT: Shimkets, Richard A.
APPLICANT: Shimkets, Richard A.
TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide Polymorphisms and
TITLE OF INVENTION: USe Thereof
FILE REFERENCE: 15966-611
FILE REFERENCE: 15966-611
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT FILING DATE: 2001-01-08
PRIOR APPLICATION NUMBER: 60/174962
PRIOR FILING DATE: 2000-01-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/755,374A CURRENT FILING DATE: 2001-01-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 28742
                              COMPUTER: HP Vectra 486/33

OPERATING SYSTEM: MSDOS version 6.2

SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:

APPLICATION NITHERED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: Accession number cg44002450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LOCATION: (25)...(26)
OTHER INFORMATION: Nucleotide deleted between bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LOCATION: (26)...(0)
OTHER INFORMATION: 1 of 2 allelic variants (13434 is other entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                             2555 ccacctcccagctttcttcttcttttttttttttctga 2590
                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                       APPLICANT: Craig A. Rosen, et. al. TITLE OF INVENTION: Human Genes, S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              44 CCAAACCCCAGATTTGTACTTTTTTTTTTTTTTCTAA 9
CLASSIFICATION:
                                                                                                                                                                                        STATE:
                                                                                                                                                                                                      STREET: 9410 Key West Avenue CITY: Rockville
                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                          ADDRESSEE:
                                                                                                                                                      20850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                    Maryland
                                                                                                                                                                      USA
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                                                                                                                                                                                                                                          Human Genome Sciences, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.6%;
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                                                                                                                                                                                                                                                                                          Human Genes, Sequences,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 23.2; DB Pred. No. 4e+05; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 23.2;
Pred. No. 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2; DB;
4e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 29;
                                                                                                                                                                                                                                                                                          and Expression Products -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
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PRIOR APPLICATION DATA: APPLICATION NUMBER:

FILING DATE:

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TOPOLOGY: lir
US-08-346-731-6670
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US-08 420-856-6670
Sequence 6670, Application US/08420856
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:

NAME: Benson, Bob

REGISTRATION NUMBER: 90,446

REFERENCE, DOCKET NUMBER: P016

TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504

TELEPAX: (301) 309-8512

INFORMATION FOR SEQ ID NO: 6670:
SEQUENCE CHARACTERISTICS:
LENGTH: 50 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
US-08-420-856-6670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.6%;
Best Local Similarity 66.7%;
Matches 32; Conservative
                                                                                      TELEFAX: (201) 994-174
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                          ATTORNEY AGENT INFORMATION:
NAME: Olstein, Elliot
REGISTRATION NUMBER: 24,025
REFERENCE/DOCKET NUMBER: 3258
TELECOMMUNICATION INFORMATION:
TELEPHONE: (201) 994-1700
TELEFAX: (201) 994-1744
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4mb storage
COMPUTER: HP Vectra 486/33
OPERATING SYSTEM: MSDOS version 5.0
SOFTWARE: ASCII Text
CURRENT APPLICATION NUMBER: US/08/420,856
APPLICATION NUMBER: US/08/420,856
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Rosen, Craig A.
APPLICANT: Ruben, Steve M.
APPLICANT: Dillon, Patrick J.
     TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
420-05-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: HU NUMBER OF SEQUENCES: 1 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                             FILING DATE:
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Roseland
STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
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                                                                                                                                                                                                                                                                           APPLICATION NUMBER:
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6 Becker Farm Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Li, Haddong
Li, Haddong
Haseltine, William A.
NVENTION: Human Genes, Sequences, and Expression Products -
SPOUENCES: 12477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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Pred. No. 4.6e+05;
0; Mismatches 16
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Query Match
Best Local Similarity
Matches 32; Conserv
2556 cacctcccagctttcttcttcttcttttttttttttttctgaaaaaaggaaaaag 2603
                                     Conservative
                                                 0.6%;
                                     Score 23; DB 8;
Pred. No. 4.6e+05;
0; Mismatches 16
                                         16;
                                                                  Length 50;
                                         Indels
                                           0;
                                           Gaps
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Search completed: September 18, Job time: 9827 sec 2002, 02:37:20 B

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Minimum
Maximum
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Perfect score:
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seq length: 50
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1: /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq:*

3: /cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:*

4: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*

5: /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:*

6: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*

7: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*

8: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:*
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Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hits satisfying chosen parameters:
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Maximum Match 100%
Listing first 45 summaries
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3713
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 September 17, 2002, 23:59:13;
       Copyright
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       attaacaggccgtggttagg.....aaactttaatgagttattta
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         GenCore version 4.5 (c) 1993 - 2000 Compugen
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US-10-181-831-5024
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US-09-396-196G-83095
US-09-396-196G-83089
US-10-131-831-77610
US-10-027-632-177610
US-10-027-632-177610
US-10-027-632-177600
US-10-131-831-4588
US-10-027-632-177600
US-10-170-097-805
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US-09-14-101-10
US-09-718-321A-50
US-10-170-097-1292
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                        Sequence 83095, A Sequence 50, Appl Sequence 50, Appl Sequence 83095, A Sequence 83089, A Sequence 177620, A Sequence 177620, Sequence 177600, Sequence 177600, Sequence 177600, Sequence 177601, Appl Sequence 7704, Appl Sequence 7704, Appl Sequence 10, Appl Sequence 10, Appl Sequence 10, Appl Sequence 1292, Appl Sequence 17768, A Sequence 1292, Appl Sequence 1292, Appl Sequence 130, Appl Sequence 50, 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Description
    Appli
Appli
       .. .. .. .. .. .. .. .. .. .. ..
APPLICANT: Wohlgemuth, Jay
APPLICANT: Fry, Kirk
APPLICANT: Fry, Kirk
APPLICANT: Ly, Ngoc
TITLE OF INVENTION: METHODS AND COMPOSITIONS FO:
TITLE OF INVENTION: TRANSPLANT REJECTION
FILE REFERENCE: 506612000121
CURRENT APPLICATION NUMBER: US/10/131,831
CURRENT APPLICATION NUMBER: US/10/06,290
PRIOR APPLICATION NUMBER: US 60/296,764
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Query Match  Best Local Similarity 96.0%; Pred. No. 5e+ Matches 24; Conservative 0; Mismatches  Qy 2110 agcccttgccccagcaatgctgaat 2134  Lillillillillillillillillilli  Db 1 agcccttgccccaggaatgctgaat 25  RESULT 2 US-10-131-831-5024/c ; Sequence 5024, Application US/10131831 ; GENERAL INFORMATION:	RESULT 1 US-09-396-196G-83096 Sequence 83096, Application US/09396196G GENERAL INFORMATION: APPLICANT: Michael Mittmann APPLICANT: David Mack APPLICANT: David Lockhart APPLICANT: David Lockhart APPLICANT: David Lockhart APPLICANT: Affymetrix, Inc. TITLE OF INVENTION: Methods of Genetic Analys FILE REFERENCE: 3101.1 CURRENT APPLICATION NUMBER: US/09/396,196G CURRENT FILING DATE: 1999-09-15 PRIOR APPLICATION NUMBER: 60/100,678 PRIOR FILING DATE: 1998-09-17 NUMBER OF SEQ ID NOS: 127806 SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 83096 LENGTH: 25 TYPE: DNA ORGANISM: mus musculus	1 396-196G-83096 396-196G-83096 BALL THEODER APPLICATION US/093	5 20 0.5 48 7 US- 5 20 0.5 48 7 US-	1 20 0.5 47 7 US-10- 2 20 0.5 48 5 US-09-	0 20 0.5 40 4 US-08-10-	7 20.2 0.5 50 6 US-10-	5 20.2 0.5 48 / US-10- 6 20.2 0.5 50 5 US-09-	33 20.2 0.5 47 7 US-10- 34 20.2 0.5 48 5 US-09-	0.5 47 5 US-09- 0.5 47 5 US-09-	29 20.2 0.5 25 5 US-09-	7 20.2 0.5 25 5 US-09- 8 20.2 0.5 25 5 US-09-	6 20.4 0.5 50 6 US-10-	
	23.4; DB 5; Length 25; No. 5e+04; smatches 1; Indels 0; Gaps 0	, 19	ALIGNMENTS 96196G	03-138B-21 Sequence 21, 50-407-21 Sequence 21, 56-306-7182 Sequence 718	397-804 Sequence 804, 289B-1326 Sequence 1326	103-1012 sequence 1012 103-245 sequence 245,	331-4502 Sequence 4503	\$06-71/2 Sequence 71/3 192-3196 Sequence 3190	70-097-1167 Sequence 1167 30-289B-1167 Sequence 1167	198A-24 Sequence	96-196G-92147 Sequence 9214	96-196G-83088 Sequence 83088, 96-196G-83097 Sequence 83097,	1-831-391 Sequence 3

FOR DIAGNOSING AND MONITORING

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RESULT 3
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                                                                                   US-10-189-360-50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR FILING DATE: 2001-06-08
NUMBER OF SEQ ID NOS: 9190
SOFTWARE: PatentIn version 3.1
SEQ ID NO 5024
LENGTH: 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 50, Application US/10189360 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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                             Query Match
Matches
              Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1214 gacttettacaagacatecetgatggtettttttttggatgetga 1257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/10/189,360
FILING DATE: 02-Unl-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/718,904
ETLING DATE: 24-SEP-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sosnowski, Barbara A.
TITLE OF INVENTION: COMPOSITIONS CONTAINING NUCLEIC ACIDS AND LIGANDS
FOR THERAPE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Baird,
                                                                                                  SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                       FEATURE:
                                                                                                                                                                     TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Seattle
STATE: Washington
                                                                                                                                                                                       TOPOLOGY: linear
                                                                                                                                                                                                                            LENGTH: 45 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                     NAME: Nottenburg Ph.D., Carol REGISTRATION NUMBER: 39,317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 6300 Columbia Center,
                                                                                                                      NAME/KEY: (Ser4Gly)2
LOCATION: 3..45
                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 760100.415C1
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Conservative
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70.5%;
              0.6%;
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            Score 22.6; DB 6; Length 45; Pred. No. 1e+05;
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Pred. No. 7.
Mismatches
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GENERAL INFORMATION:
APPLICANT: Michael Mittmann
APPLICANT: Michael Mittmann
APPLICANT: David Mack
APPLICANT: David Lockhart
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3101.1
CURRENT APPLICATION NUMBER: US/09/396,196G
CURRENT FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: 60/100,678
PRIOR FILING DATE: 1998-09-17
NUMBER OF SEQ ID NOS: 127806
SOSTWARE: PASTSEQ for Windows Version 4.0
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US-09-396-196G-83095
; Sequence 83095, Application US/09396196G
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                                                                                                                                                                          ; ORGANISM: Homo sapiens US-10-131-831-7656
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                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: US 10/006,290
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
NUMBER OF SEQ ID NOS: 9190
SOFTWARE: Patentin version 3.1
SEQ ID NO 7656
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LENGTH: 25
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Best Local (
                                                                         Query Match
Best Local Similarity
Matches 25; Conserv
                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Ly, NGOC
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING
TITLE OF INVENTION: TRANSPLANT REJECTION
FILE REFERENCE: 506612000121
CURRENT APPLICATION NUMBER: US/10/131,831
CURRENT FILING DATE: 2002-08-05
CURRENT FILING DATE: 2002-08-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Wohlgemuth, Jay
APPLICANT: Fry, Kirk
APPLICANT: Woodward, Robe
APPLICANT: Ly, Ngoc
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TYPE: DNA
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                      1638 gagacactgtctcagtgcatcgcccaggct 1667
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48 GAGACAGGGTCTCACTGTATTGCCCCAGGCT 19
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Woodward, Robert
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Pred. No. 8.9e+04;
D; Mismatches 1
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Pred. No. 1
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1.5e+05
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RESULT

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; TYPE: DNA
; ORGANISM: mus musculus
US-09-396-196G-83082
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; ORGANISM: mus musculus
US-09-396-196G-83089
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Best Local Similarity 92.0%;
Matches 23; Conservation
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Best Local Similarity
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APPLICANT: Wohlgemuth, Jay
APPLICANT: Wohlgemuth, Jay
APPLICANT: Woodward, Robert
APPLICANT: Woodward, Robert
APPLICANT: Ly, Ngoc
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING
TITLE OF INVENTION: TRANSPLANT REJECTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/396,196G
CURRENT FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: 60/100,678
PRIOR FILING DATE: 1998-09-17
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TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3101.1
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APPLICANT: David Lockhart
APPLICANT: Affymetrix, In
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TITLE OF INVENTION: Methods of Genetic Analysis
FILE REFERENCE: 3101.1
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Pred. No. 1.3e+05;
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Pred. No. 1.3e+05;
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RESULT 10
US-10-027-632-177620/c
US-10-027-632-177620/c
; Sequence 177620, Application US/10027632
; GENERAL INFORMATION:
; APPLICANY: Wang, David G.
; TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TYPE: DNA; Homo sapiens; ORGANISM: Homo sapiens US-10-131-831-4391
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CURRENT APPLICATION NUMBER: US/10/131,831
CURRENT FILING DATE: 2002-08-05
PRIOR APPLICATION NUMBER: US 10/006,290
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
PRIOR FILING DATE: 2001-06-08
NUMBER OF SEQ ID NOS: 9190
SOFTWARE: PatentIn version 3.1
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Best Local Similarity
Matches 28; Conserv
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LENGTH: 50
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CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
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PRIOR
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PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
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                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
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Pred. No. 1
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Pred. No. 2e+05;
0; Mismatches
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REFERENCE:

OF INVENTION:

ION: Polymorphisms in the Human Genome 108827.129

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APPLICANT: LY, NGOC

TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING AND MONITORING
FILE OF INVENTION: TRANSPLANT REJECTION
FILE REFERENCE: 506612000121
CURRENT APPLICATION NUMBER: US/10/131,831
CURRENT FILING DATE: 2002-08-05
PRIOR APPLICATION NUMBER: US 10/006,290
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR APPLICATION NUMBER: US 60/296,764
PRIOR FILING DATE: 2001-06-08
NUMBER: DATE: 2001-06-08
NUMBER: PATENTING DATE: 2001-06-08
NUMBER: DATE: 2001-06-08
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RESULT 12
US-10;055-728-32
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CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-04-20
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PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 60/146,002 PRIOR FILING DATE: 1999-08-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
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                                                                                                                                   3247 acaagagttgtggttttccatctgatccttc 3277
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Local Similarity 71.8%;
nes 28; Conservativo
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                                                                                                      17 acataagttatggttttccatcttatctgtc 47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Application US/10131831
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Pred. No. 2
                                                                                                                                                                                                             Mismatches
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2939 ggttgtttttctaaatgtgaagacttaccaaatgaattt 2977

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; :OTHER INFORMATION: oligo (dT) primer with a 5' US-10-055-728-32
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NUMBER OF SEQ ID NOS: 156
SOFTWARE: PatentIn version 3.1
SEQ ID NO 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Seguence 32, Application US/10055728 GENERAL INFORMATION:
whery Match
Elest Local Similarity by...
The conservative
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                                                                                                                             ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GURRENT APPLICATION NUMBER: US/10/055,728
GURRENT FILING DATE: 2002-04-19
ERIOR APPLICATION NUMBER: 60/325,722
ERIOR FILING DATE: 2001-09-28
ERIOR APPLICATION NUMBER: EP 0120373.2
ERIOR APPLICATION NUMBER: EP 0120373.2
ERIOR FILING DATE: 2001-09-28
ERIOR APPLICATION NUMBER: EP 01200228.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Cornelissen, Marion
TITLE OF INVENTION: MEANS AND METHODS FOR TREATMENT EVALUATION
EVILE REFERENCE: 5244US (REN/P551900500)
                                                                                                                                                                                                          SEQ ID NO 177600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MPPLICANT: van der Kuyl,
                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR FILING DATE: 2000-02-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            凾-027-632-177600/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RIOR APPLICATION NUMBER: US 60/218,006 RIOR FILING DATE: 2000-07-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-08-28
PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
PRIOR FILING DATE: 1999-08-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             URRENT APPLICATION NUMBER: US/10/027,632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TILE OF INVENTION: Identification and Mapping of Single ITLE OF INVENTION: Polymorphisms in the Human Genome ILE REFERENCE: 108827.129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PPLICANT: Wang, Da
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     quence 177600, Application US/10027632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
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                                                                                                                                                                                         CENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                               for Windows Version 4.0
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                               1; Mismatches
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Pred. No. 2.3e+05;
                                                  Score 21; DB 7;
Pred. No. 2.5e+05;
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                                                                    Length 43;
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                                   Gaps
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Вþ

39 GCTTTTTGTTATGAGTGTGAACRTTTATCACTTGAATTT 1

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TITLE OF INVENTION: BIALLELIC MARKERS DERIVED FROM GENOMIC REGIONS CARRYING TITLE OF INVENTION: GENES INVOLVED IN ARACHIDONIC ACID METABOLISM FILE REFERENCE: GEN-T1114XC2DI .

CURRENT APPLICATION NUMBER: US/10/170,097

CURRENT FILING DATE: 2002-06-10

PRIOR APPLICATION NUMBER: US 09/641,638

PRIOR FILING DATE: 2000-08-16

PRIOR APPLICATION NUMBER: US 09/502,330

PRIOR FILING DATE: 2000-02-11

PRIOR APPLICATION NUMBER: US 60/133,200

PRIOR FILING DATE: 1999-05-07

PRIOR FILING DATE: 1999-05-07

PRIOR APPLICATION NUMBER: US 60/133,200

PRIOR FILING DATE: 1999-05-07

PRIOR APPLICATION NUMBER: US 60/139,207

PRIOR APPLICATION NUMBER: US 60/139,207

PRIOR APPLICATION NUMBER: US 60/119,917

PRIOR PRIOR FILING DATE: 1999-02-12
                                                                                                                                                                                TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:
US-10-156-306-7691
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: 12-740-118 : polymorphic base C or US-10-170-097-805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-10-170-097-805
 Query Match 0.6%;
Best Local Similarity 66.7%;
Matches 30; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.6%;
Best Local Similarity 65.9%;
Matches 27; Conservative
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SOFTWARE: Patent.pm
SEQ ID NO 805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Blumenfeld, Marta
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LOCATION: 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bougueleret, Lydie
Chumakov, Ilya
 Score 21; DB 7; Le
Pred. No. 2.6e+05;
0; Mismatches 15;
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Pred. No. 2.6e+05;
2; Mismatches 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               12; Indels
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                                      Length 48;
   Indels
                                                                                                               Enzymatic Nucleic Acid
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Gaps
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Search completed: September 18, 2002, 02:48:37 Job time: 10164 sec